

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/538,885
Source: PCT
Date Processed by STIC: 06/22/2005

ENTERED



PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/538,885

DATE: 06/22/2005

TIME: 10:20:59

Input Set : A:\Sequence Listing US.ST25.txt
 Output Set: N:\CRF4\06222005\J538885.raw

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3 <110> APPLICANT: Nessler, Craig
4     Lorence, Argelia
5     Chevone, Boris
6     Mendes, Pedro
8 <120> TITLE OF INVENTION: Manipulation of Ascorbic acid Levels in Plants
10 <130> FILE REFERENCE: 01640421aa
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/538,885
C--> 12 <141> CURRENT FILING DATE: 2005-06-14
12 <150> PRIOR APPLICATION NUMBER: US 60/433,787
13 <151> PRIOR FILING DATE: 2002-12-17
15 <150> PRIOR APPLICATION NUMBER: PCT/US03/27779
16 <151> PRIOR FILING DATE: 2003-09-08
18 <160> NUMBER OF SEQ ID NOS: 4
20 <170> SOFTWARE: PatentIn version 3.2
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 21
24 <212> TYPE: DNA
25 <213> ORGANISM: Artificial
27 <220> FEATURE:
28 <223> OTHER INFORMATION: Synthetic oligonucleotide primer for miox gene (forward)
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31 cccatggcga tctctgttga g                                21
34 <210> SEQ ID NO: 2
35 <211> LENGTH: 21
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial
39 <220> FEATURE:
40 <223> OTHER INFORMATION: Synthetic oligonucleotide primer for miox gene (reverse)
42 <400> SEQUENCE: 2
43 ccggatccctc accacacctaa g                                21
46 <210> SEQ ID NO: 3
47 <211> LENGTH: 954
48 <212> TYPE: DNA
49 <213> ORGANISM: Arabidopsis thaliana
52 <220> FEATURE:
53 <221> NAME/KEY: Source
54 <222> LOCATION: (1)..(954)
55 <223> OTHER INFORMATION: nucleic acid sequence of A. thaliana miox4 cDNA PCR product
57 <400> SEQUENCE: 3
58 atgacgatct ctgttgagaa gccgattttt gaagaagagg tttctgcatt cgagaagagt      60
60 gggacaata tcggagagtt gaaattggac ggaggattt cgatgccgaa aatggacacc      120
62 aatgacgacg aagttttt ggctccttag atgaatgcatttggccgcca attcaggac      180
64 tacgatgtt agagtgagag gcaaaaggc gtcgaagagt tttacagatt acgacacatc      240

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66	aaccaaactg	tcgactttgt	aaaaagatg	aggctgaat	atggaaaatt	agataaaatg	300									
68	gtgatgagca	tttggaaatg	ttgtgagctt	ctcaatgagg	ttgtggatga	gagtgatcca	360									
70	gatcttgacg	agccccagat	tcagcatttg	cttcaatctg	ccgaagccat	ccgcaaagat	420									
72	taccctaatg	aagattggct	tcatctgacc	gctcttatcc	atgatcttg	gaaagttatt	480									
74	actcttccac	aattcgagg	acctcctcaa	tggctgttg	ttggtgacac	attccctgtt	540									
76	ggatgtgcac	ttgatgaatc	taacgtacat	cacaagtact	ttgtggaaaa	cccagattt	600									
78	cacaacgaaa	cctacaacac	taaaaatggg	atttactctg	.aagggtgtgg	attaaacaat	660									
80	gtcatgatgt	cttggggcca	tgacgactac	atgtacctgg	tggctaaaga	aaacggaagt	720									
82	accttgcgt	cggctggaca	gtttatcata	agataccact	ccttttaccc	tttgcacacg	780									
84	gctggagaat	acacccatct	tatgaacgag	gaagacaagg	agaatctgaa	gtggctacac	840									
86	gttttcaaca	agtacgactt	gtacagtaag	agcaaagttc	acgttgatgt	ggagaaggtc	900									
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93	<212>	TYPE:	PRT													
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97	<220>	FEATURE:														
98	<221>	NAME/KEY:	Source													
99	<222>	LOCATION:	(1)..(318)													
100	<223>	OTHER INFORMATION:	Amino acid sequence of A. thaliana miox4 cDNA product													
102	<400>	SEQUENCE:	4													
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105	1				5			10				15				
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109						20			25			30				
112	Phe	Ser	Met	Pro	Lys	Met	Asp	Thr	Asn	Asp	Asp	Glu	Ala	Phe	Leu	Ala
113						35			40			45				
116	Pro	Glu	Met	Asn	Ala	Phe	Gly	Arg	Gln	Phe	Arg	Asp	Tyr	Asp	Val	Glu
117						50			55			60				
120	Ser	Glu	Arg	Gln	Lys	Gly	Val	Glu	Glu	Phe	Tyr	Arg	Leu	Arg	His	Ile
121						65			70			75			80	
124	Asn	Gln	Thr	Val	Asp	Phe	Val	Lys	Lys	Met	Arg	Ala	Glu	Tyr	Gly	Lys
125						85			90			95				
128	Leu	Asp	Lys	Met	Val	Met	Ser	Ile	Trp	Glu	Cys	Cys	Glu	Leu	Leu	Asn
129						100			105			110				
132	Glu	Val	Val	Asp	Glu	Ser	Asp	Pro	Asp	Leu	Asp	Glu	Pro	Gln	Ile	Gln
133						115			120			125				
136	His	Leu	Leu	Gln	Ser	Ala	Glu	Ala	Ile	Arg	Lys	Asp	Tyr	Pro	Asn	Glu
137						130			135			140				
140	Asp	Trp	Leu	His	Leu	Thr	Ala	Leu	Ile	His	Asp	Leu	Gly	Lys	Val	Ile
141	145					145			150			155			160	
144	Thr	Leu	Pro	Gln	Phe	Gly	Gly	Leu	Pro	Gln	Trp	Ala	Val	Val	Gly	Asp
145						165			170			175				
148	Thr	Phe	Pro	Val	Gly	Cys	Ala	Phe	Asp	Glu	Ser	Asn	Val	His	His	Lys
149						180			185			190				
152	Tyr	Phe	Val	Glu	Asn	Pro	Asp	Phe	His	Asn	Glu	Thr	Tyr	Asn	Thr	Lys
153						195			200			205				
156	Asn	Gly	Ile	Tyr	Ser	Glu	Gly	Cys	Gly	Leu	Asn	Asn	Val	Met	Met	Ser
157						210			215			220				

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160 Trp Gly His Asp Asp Tyr Met Tyr Leu Val Ala Lys Glu Asn Gly Ser
161 225 230 235 240
164 Thr Leu Pro Ser Ala Gly Gln Phe Ile Ile Arg Tyr His Ser Phe Tyr
165 245 250 255
168 Pro Leu His Thr Ala Gly Glu Tyr Thr His Leu Met Asn Glu Glu Asp
169 260 265 270
172 Lys Glu Asn Leu Lys Trp Leu His Val Phe Asn Lys Tyr Asp Leu Tyr
173 275 280 285
176 Ser Lys Ser Lys Val His Val Asp Val Glu Lys Val Glu Pro Tyr Tyr
177 290 295 300
180 Met Ser Leu Ile Lys Lys Tyr Phe Pro Glu Asn Leu Arg Trp
181 305 310 315

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/22/2005
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2

VERIFICATION SUMMARY

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Input Set : A:\Sequence Listing US.ST25.txt
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L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date